



SEQUENCE LISTING

<110> Pharmacia AB
<120> Protein Cluster II
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<140> US/10/029,359
<141> 2001-12-21
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<170> PatentIn version 3.0
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Lys Ser Trp Leu Asn Phe Leu Thr Phe Leu Tyr Gly Ser Ala Ile Gly
15 20 25

ttt att tta ttt tct cag cta ctt agt att ttg ttg gga gaa gag ggt 149
Phe Ile Leu Phe Ser Gln Leu Leu Ser Ile Leu Leu Gly Glu Glu Gly
30 35 40

gac acc cag act aat gtt ctt cat aat gat cct cat gcg agg cat tca 197
Asp Thr Gln Thr Asn Val Leu His Asn Asp Pro His Ala Arg His Ser
45 50 55

gat gat aat gga cag aat cat cta gga gga caa atg aac ttc aat gca 245
Asp Asp Asn Gly Gln Asn His Leu Gly Gly Gln Met Asn Phe Asn Ala
60 65 70 75

gat tct agc caa cgt aaa gat gag aac aca gaa atc gct gaa aac ctc 293
Asp Ser Ser Gln Arg Lys Asp Glu Asn Thr Glu Ile Ala Glu Asn Leu
80 85 90

tat nag caa gtt aaa att ctt tgc tgg gtt atg aca ggc tct caa aac 341
Tyr Xaa Gln Val Lys Ile Leu Cys Trp Val Met Thr Gly Ser Gln Asn
95 100 105

cta cag aaa aag gcc aaa cat gtc aaa gct aca tgg gcc cag cgt tgt 389
Leu Gln Lys Lys Ala Lys His Val Lys Ala Thr Trp Ala Gln Arg Cys
110 115 120

cta aaa gta ttt ttt atg agt tca gaa gaa aat aaa gac ttc cgt gct 437
Leu Lys Val Phe Phe Met Ser Ser Glu Glu Asn Lys Asp Phe Arg Ala

125	130	135	
gtg gga ttg aaa acc aaa gca ggc aga gat gag cta tac tgg aaa aca			485
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Gln Leu Leu Ser Ile Leu Leu Gly Glu Glu Gly Asp Thr Gln Thr Asn
35 40 45

Val Leu His Asn Asp Pro His Ala Arg His Ser Asp Asp Asn Gly Gln
50 55 60

Asn His Leu Gly Gly Gln Met Asn Phe Asn Ala Asp Ser Ser Gln Arg
65 70 75 80

Lys Asp Glu Asn Thr Glu Ile Ala Glu Asn Leu Tyr Xaa Gln Val Lys
85 90 95

Ile Leu Cys Trp Val Met Thr Gly Ser Gln Asn Leu Gln Lys Lys Ala
100 105 110

Lys His Val Lys Ala Thr Trp Ala Gln Arg Cys Leu Lys Val Phe Phe
115 120 125

Met Ser Ser Glu Glu Asn Lys Asp Phe Arg Ala Val Gly Leu Lys Thr
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Lys Ala Gly Arg Asp Glu Leu Tyr Trp Lys Thr Ile Asn Leu Phe
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Thr Gly Val Thr Asp Lys Leu Tyr Gln Lys Met Lys Ile Leu Cys Trp
5 10 15

att atg aca gga cct caa aat cta gaa aaa aag atc aga cgc atc aga 153
Ile Met Thr Gly Pro Gln Asn Leu Glu Lys Lys Ile Arg Arg Ile Arg
20 25 30

gat aca tgg gcc cag ggt tgc aat aaa gcg ttg ttt atg agc tca aaa 201
Asp Thr Trp Ala Gln Gly Cys Asn Lys Ala Leu Phe Met Ser Ser Lys
35 40 45

gaa aat aaa gac ttc tct act gtg gga tta cac acc aaa gaa gac aga 249
Glu Asn Lys Asp Phe Ser Thr Val Gly Leu His Thr Lys Glu Asp Arg
50 55 60 65

aac caa ctg tcc tgg aaa ata gtt aaa gct ttt cta tat gct cat gac 297
Asn Gln Leu Ser Trp Lys Ile Val Lys Ala Phe Leu Tyr Ala His Asp
70 75 80

cat tat ctg gaa tac atg gat tgg ttc atg aaa gca gat gat gat ata 345
His Tyr Leu Glu Tyr Met Asp Trp Phe Met Lys Ala Asp Asp Asp Ile
85 90 95

tgt ata tat atc aca ttg gac aac ttg aaa tgg ctt ctc aca aac tat 393
Cys Ile Tyr Ile Thr Leu Asp Asn Leu Lys Trp Leu Leu Thr Asn Tyr
100 105 110

aac cct gat gaa tcc act tac ttt ggg aaa aga ttt aag cac tgc aga 441
Asn Pro Asp Glu Ser Thr Tyr Phe Gly Lys Arg Phe Lys His Cys Arg
115 120 125

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 35 40 45

Lys Glu Asn Lys Asp Phe Ser Thr Val Gly Leu His Thr Lys Glu Asp
 50 55 60

Arg Asn Gln Leu Ser Trp Lys Ile Val Lys Ala Phe Leu Tyr Ala His
 65 70 75 80

Asp His Tyr Leu Glu Tyr Met Asp Trp Phe Met Lys Ala Asp Asp Asp
 85 90 95

Ile Cys Ile Tyr Ile Thr Leu Asp Asn Leu Lys Trp Leu Leu Thr Asn
 100 105 110

Tyr Asn Pro Asp Glu Ser Thr Tyr Phe Gly Lys Arg Phe Lys His Cys
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Arg Lys Gln Asp Tyr Met Thr Gly Gly Ala Gly Tyr Val Leu Ser Lys
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Glu
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tct agc caa cat aaa gat gag aac aca gac att gct gaa aac ctc tat 97
 Ser Ser Gln His Lys Asp Glu Asn Thr Asp Ile Ala Glu Asn Leu Tyr
 20 25 30

cag aaa gtt aga att ctt tgc tgg gtt atg acc ggc cct caa aac cta 145
 Gln Lys Val Arg Ile Leu Cys Trp Val Met Thr Gly Pro Gln Asn Leu
 35 40 45

gag aaa aag gcc aaa cac gtc aaa gct act tgg gcc cag cgt tgt aac 193
 Glu Lys Lys Ala Lys His Val Lys Ala Thr Trp Ala Gln Arg Cys Asn
 50 55 60

aaa gtg ttg ttt atg agt tca gaa gaa aat aaa gac ttc cct gct gtg 241
 Lys Val Leu Phe Met Ser Ser Glu Glu Asn Lys Asp Phe Pro Ala Val
 65 70 75 80

gga ctg aaa acc aaa gaa ggc aga gat caa cta tac tgg aaa aca att 289
 Gly Leu Lys Thr Lys Glu Gly Arg Asp Gln Leu Tyr Trp Lys Thr Ile
 85 90 95

aaa gct ttt cag tat gtt cat gaa cat tat tta caa gat gct gat tgg Lys Ala Phe Gln Tyr Val His Glu His Tyr Leu Gln Asp Ala Asp Trp 100 105 110	337
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tat gta cta agc aaa gaa gcc ttg aaa aga ttt gtt gat gca ttt aaa Tyr Val Leu Ser Lys Glu Ala Leu Lys Arg Phe Val Asp Ala Phe Lys 165 170 175	529
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aga tgc atg gaa att atg aat gta gaa gca gga gat tcc aga gat acc Arg Cys Met Glu Ile Met Asn Val Glu Ala Gly Asp Ser Arg Asp Thr 195 200 205	625
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aaa ggt tat cta cct aga acg ttt tgg tac tgg aat tac aac tat tat Lys Gly Tyr Leu Pro Arg Thr Phe Trp Tyr Trp Asn Tyr Asn Tyr Tyr 225 230 235 240	721
cct cct gta gag ggt cct ggt tgc tgc tct gat ctt gca gtt tct ttt Pro Pro Val Glu Gly Pro Gly Cys Cys Ser Asp Leu Ala Val Ser Phe 245 250 255	769
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gaa cgt ata cta aag gaa att agt caa gca aac aaa aat gaa gat aca Glu Arg Ile Leu Lys Glu Ile Ser Gln Ala Asn Lys Asn Glu Asp Thr 290 295 300	913
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tatgtctata tatatgagga acttgtgttt tttaaagtgt ggccaggtag aggaactaga	1204

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Gln Lys Val Arg Ile Leu Cys Trp Val Met Thr Gly Pro Gln Asn Leu
 35 40 45

Glu Lys Lys Ala Lys His Val Lys Ala Thr Trp Ala Gln Arg Cys Asn
 50 55 60

Lys Val Leu Phe Met Ser Ser Glu Glu Asn Lys Asp Phe Pro Ala Val
 65 70 75 80

Gly Leu Lys Thr Lys Glu Gly Arg Asp Gln Leu Tyr Trp Lys Thr Ile
 85 90 95

Lys Ala Phe Gln Tyr Val His Glu His Tyr Leu Gln Asp Ala Asp Trp
 100 105 110

Phe Leu Lys Ala Asp Asp Asp Thr Tyr Val Ile Leu Asp Asn Leu Arg
 115 120 125

Trp Leu Leu Ser Lys Tyr Asp Pro Glu Glu Pro Ile Tyr Phe Gly Arg
 130 135 140

Arg Phe Lys Pro Tyr Val Lys Gln Gly Tyr Met Ser Gly Gly Ala Gly
 145 150 155 160

Tyr Val Leu Ser Glu Ala Leu Lys Arg Phe Val Asp Ala Phe Lys
 165 170 175

Thr Asp Thr His Ser Ser Ser Ile Glu Asp Leu Ala Leu Gly
 180 185 190

Arg Cys Met Glu Ile Met Asn Val Glu Ala Gly Asp Ser Arg Asp Thr
195 200 205

Ile Gly Lys Glu Thr Phe His Pro Phe Val Pro Glu His His Leu Ile
210 215 220

Lys Gly Tyr Leu Pro Arg Thr Phe Trp Tyr Trp Asn Tyr Asn Tyr Tyr
225 230 235 240

Pro Pro Val Glu Gly Pro Gly Cys Cys Ser Asp Leu Ala Val Ser Phe
245 250 255

His Tyr Val Asp Ser Thr Thr Met Tyr Glu Leu Glu Tyr Leu Val Tyr
260 265 270

His Leu Arg Pro Tyr Gly Tyr Leu Tyr Arg Tyr Gln Pro Thr Leu Pro
275 280 285

Glu Arg Ile Leu Lys Glu Ile Ser Gln Ala Asn Lys Asn Glu Asp Thr
290 295 300

Lys Val Lys Leu Gly Asn Pro
305 310